RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/581,183

Source: IFWP

Date Processed by STIC: 6/19/06

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IFWP

RAW SEQUENCE LISTING   DATE: 06/19/2006
PATENT APPLICATION: US/10/581,183   TIME: 11:23:11

Input Set : A:\14875-162US1.txt
Output Set: N:\CRF4\06192006\J581183.raw

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   3 <110> APPLICANT: Tsunoda, Hiroyuki
   4        Habu, Kiyoshi
   6 <120> TITLE OF INVENTION: EXPRESSION SYSTEMS USING MAMMALIAN BETA-ACTIN
PROMOTER
   8 <130> FILE REFERENCE: 14875-162US1
C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/581,183
C--> 10 <141> CURRENT FILING DATE: 2006-06-01
  10 <150> PRIOR APPLICATION NUMBER: PCT/JP2004/018006
  11 <151> PRIOR FILING DATE: 2004-12-03
  13 <150> PRIOR APPLICATION NUMBER: JP 2003-405269
  14 <151> PRIOR FILING DATE: 2003-12-03
  16 <160> NUMBER OF SEQ ID NOS: 39
  18 <170> SOFTWARE: PatentIn version 3.1
  20 <210> SEQ ID NO: 1
  21 <211> LENGTH: 1577
  22 <212> TYPE: DNA
  23 <213> ORGANISM: Mus musculus
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  30 acatccacac ccagagggtc ctggggtggt tgggtgaccc ccagaatgca ggcctagtaa     180
  32 ccgagacatt gaatggggca gtgtccacaa gggcggaggc tattcctgta catctgggcc     240
  34 tacggagcca gcacccatcg ccaaaactct tcatcctctt cctcaatctc gctttctctc     300
  36 tcgctttttt tttttttcct tcttcttttt ttttttttt ttcaaaagga ggggagaggg     360
  38 ggtaaaaaaa tgctgcactg tgcggcgagg ccggtgagtg agcgacgcgg agccaatcag     420
  40 cgcccgccgt tccgaaagtt gccttttatg gctcgagtgg ccgctgtggc gtcctataaa     480
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  44 tgcagctcct tcgttgccgg tccacacccg ccaccaggta agcagggacg ccgggcccag     600
  46 cgggccttcg ctctctcgtg gctagtacct cactgcaggg tcctgaggat cactcagaac     660
  48 ggacaccatg ggcgggtgga gggtggtgcc gggccgcgga gcggacactg gcacagccaa     720
  50 ctttacgcct agcgtgtaga ctctttgcag ccacattccc gcggtgtaga cactcgtggg     780
  52 cccgctcccg ctcggtgcgt ggggcttggg gacacactag ggtcgcggtg gggcatttg     840
  54 atgagccggt gcggcttgcg ggtgttaaaa gccgtattag gtccatcttg agagtacaca     900
  56 gtattgggaa ccagacgcta cgatcacgcc tcaatggcct ctgggtcttt gtccaaaccg     960
  58 gtttgcctat tcggcttgcc gggcgggcgg gcgggcgggc gggcgcggca gggccggctc    1020
  60 ggccgggtgg gggctgggat gccactgcgc gtgcgctctc tatcactggg catcgaggcg    1080
  62 cgtgtgcgct agggagggag ctcttcctct cccctcttc ctagttagct gcgcgtgcgt    1140
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RAW SEQUENCE LISTING   DATE: 06/19/2006
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 91 acatccacac ccagagggtc ctggggtggt tgggtgaccc ccagaatgca ggcctagtaa   180
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 99 ggtaaaaaaa tgctgcactg tgcggcgagg ccggtgagtg agcgacgcgg agccaatcag   420
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123 cgtgtgcgct agggagggag ctcttcctct cccctcttc ctagttagct gcgcgtgcgt  1140
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133 gtaataatgc ggccggtctg cgcttccttt gtccctgag cttgggcgcg cgcccctgg  1440
135 cggctcgagc ccgcggcttg ccggaagtgg gcagggcggc agcggctgct cttggcggcc  1500
137 ccgaggtgac tatagccttc ttttgtgtct tgatagttcg cc                    1542
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141 <211> LENGTH: 604
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143 <213> ORGANISM: Woodchuck hepatitis virus
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148 gttgctcctt ttacgctatg tggatacgct gctttaatgc ctttgtatca tgctattgct   120
150 tcccgtatgg ctttcatttt ctcctccttg tataaatcct ggttgctgtc tctttatgag   180
152 gagttgtggc ccgttgtcag caacgtggc gtggtgtgca ctgtgtttgc tgacgcaacc   240
154 cccactggtt ggggcattgc caccacctgt cagctccttt ccgggacttt cgctttcccc   300
156 ctccctattg ccacggcgga actcatcgcc gcctgccttg cccgctgctg gacaggggct   360
158 cggctgttgg gcactgacaa ttccgtggtg ttgtcgggga agctgacgtc ctttccatgg   420
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162 gccctcaatc cagcggacct tccttcccgc ggcctgctgc cggctctgcg gcctcttccg   540
164 cgtcttcgcc ttcgccctca gacgagtcgg atctcccttt gggccgcctc cccgcctgtc   600
166 taga                                                              604
169 <210> SEQ ID NO: 4
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172 <213> ORGANISM: Homo sapiens
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179 gacgtcaata atgacgtatg ttcccatagt aacgccaata gggactttcc attgacgtca     180
181 atgggtggag tatttacggt aaactgccca cttggcagta catcaagtgt atcatatgcc     240
183 aagtacgccc cctattgacg tcaatgacgg taaatggccc gcctggcatt atgcccagta     300
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187 catggt                                                                366
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193 <213> ORGANISM: Mus musculus
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200 gtatgatccc actatagagg actcctaccg gaaacaggtg gtcattgatg gggagacatg     180
202 tctactggac atcttagaca cagcaggtca agaagagtat agtgccatgc gggaccagta     240
204 catgcgcaca ggggagggct tcctctgtgt atttgccatc aacaacacca agtccttcga     300
206 ggacatccat cagtacaggg agcagatcaa gcgggtgaaa gattcagatg atgtgccaat     360
208 ggtgctggtg ggcaacaagt gtgacctggc tgctcgcact gttgagtctc ggcaggccca     420
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212 cgtggaggat gccttctata cactagtccg tgagattcgg cagcataaat tgcggaaact     540
214 gaacccaccc gatgagagtg gtcctggctg catgagctgc aaatgtgtgc tgtcctgaca     600
216 ccaggtgagg cagggaccag cgagacgtct ggggcagtga cctcagctag ccagatgaac     660
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221 <212> TYPE: DNA
222 <213> ORGANISM: Mus musculus
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227 ctgaccatcc agctgatcca gaaccacttt gtggacgagt atgatcccac tatagaggac     120
229 tcctaccgga acaggtggt cattgatggg gagacatgtc tactggacat cttagacaca     180
231 gcaggtcaag aagagtatag tgccatgcgg gaccagtaca tgcgcacagg ggagggcttc     240
233 ctctgtgtat ttgccatcaa caacaccaag tccttcgagg acatccatca gtacagggag     300
235 cagatcaagc gggtgaaaga ttcagatgat gtgccaatgg tgctggtggg caacaagtgt     360
237 gacctggctg ctcgcactgt tgagtctcgg caggcccagg accttgctcg cagctatggc     420
239 atcccctaca ttgaaacatc agccaagacc cggcagggcg tggaggatgc cttctataca     480
241 ctagtccgtg agattcggca gcataaattg cggaaactga acccacccga tgagagtggt     540
243 cctggctgca tgagctgcaa atgtgtgctg tcctga                               576
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247 <211> LENGTH: 189
248 <212> TYPE: PRT
249 <213> ORGANISM: Mus musculus
251 <400> SEQUENCE: 7
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258 Asp Pro Thr Ile Glu Asp Ser Tyr Arg Lys Gln Val Val Ile Asp Gly
259               35                40                45
261 Glu Thr Cys Leu Leu Asp Ile Leu Asp Thr Ala Gly Gln Glu Glu Tyr
262      50                55                60
264 Ser Ala Met Arg Asp Gln Tyr Met Arg Thr Gly Glu Gly Phe Leu Cys
265 65               70                75                80
267 Val Phe Ala Ile Asn Asn Thr Lys Ser Phe Glu Asp Ile His Gln Tyr
268               85                90                95
270 Arg Glu Gln Ile Lys Arg Val Lys Asp Ser Asp Asp Val Pro Met Val
271               100               105               110
273 Leu Val Gly Asn Lys Cys Asp Leu Ala Ala Arg Thr Val Glu Ser Arg
274          115               120               125
276 Gln Ala Gln Asp Leu Ala Arg Ser Tyr Gly Ile Pro Tyr Ile Glu Thr
277      130               135               140
279 Ser Ala Lys Thr Arg Gln Gly Val Glu Asp Ala Phe Tyr Thr Leu Val
280 145               150               155               160
282 Arg Glu Ile Arg Gln His Lys Leu Arg Lys Leu Asn Pro Pro Asp Glu
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285 Ser Gly Pro Gly Cys Met Ser Cys Lys Cys Val Leu Ser
286               180               185
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290 <211> LENGTH: 188
291 <212> TYPE: PRT
292 <213> ORGANISM: Homo sapiens
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299               20                25                30
301 Asp Pro Thr Ile Glu Asp Ser Tyr Arg Lys Gln Val Val Ile Asp Gly
302               35                40                45
304 Glu Thr Cys Leu Leu Asp Ile Leu Asp Thr Ala Gly Gln Glu Glu Tyr
305      50                55                60
307 Ser Ala Met Arg Asp Gln Tyr Met Arg Thr Gly Glu Gly Phe Leu Cys
308 65               70                75                80
310 Val Phe Ala Ile Asn Asn Thr Lys Ser Phe Glu Asp Ile His His Tyr
311               85                90                95
313 Arg Glu Gln Ile Lys Arg Val Lys Asp Ser Glu Asp Val Pro Met Val
314               100               105               110
316 Leu Val Gly Asn Lys Cys Asp Leu Pro Ser Arg Thr Val Asp Thr Lys
317          115               120               125
319 Gln Ala Gln Asp Leu Ala Arg Ser Tyr Gly Ile Pro Phe Ile Glu Thr
320      130               135               140
322 Ser Ala Lys Thr Arg Gln Gly Val Asp Asp Ala Phe Tyr Thr Leu Val
323 145               150               155               160
325 Arg Glu Ile Arg Lys His Lys Glu Lys Met Ser Lys Asp Gly Lys Lys
326               165               170               175
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398 <211> LENGTH: 50

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 06/19/2006
PATENT APPLICATION: US/10/581,183   TIME: 11:23:13

Input Set : **A:\14875-162US1.txt**
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Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:9,10,11,12,13,14,15,16,17,18,19,20,21,22,23,24,25,26,27,28,29,30,31,32,33
Seq#:34,35,36,37,38,39

VERIFICATION SUMMARY   DATE: 06/19/2006
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L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date